

POST-DOCTORAL RESEARCHER · BIOINFORMATICS & TRAINING

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Education

Doctor in science Lyon, France

INSTITUT NATIONAL DES SCIENCES APPLIQUÉES

Oct. 2011 - Nov. 2014

- In silico experimental evolution
- · Comparative genomics with phylogeny

Master degree Lyon, France

ECOLE NORMALE SUPÉRIEURE & UNIVERSITY CLAUDE BERNARD

Oct. 2010 - Oct. 2011

- Theoretical computer science
- · Complex system modelling

Master of Engineering Lyon, France

INSTITUT NATIONAL DES SCIENCES APPLIQUÉES

Oct. 2018 - Oct. 2011

· Bioinformatics and Modelling

Skills

Bioinformatics HTS data analysis (RNA-Seq, metagenomics), comparative microbial genomics, phylogenetics

Software development Python, R

Training Data analysis, Galaxy, Software development, Software Carpentry instructor

Community Galaxy Traing community manager, Mozilla Open Leader, Software Carpentry mentor

Languages French (native), English, German (B2)

Experience

Post-doctoral researcher Freiburg, Germany

GALAXY TEAM, BACKOFEN LAB, UNIVERSITY OF FREIBURG

Aug. 2016 - Today

- Data analysis of High-Throughput sequencing data (RNA-seq, metagenomics, etc)
- Training on Galaxy use, data analysis and software development
- Development and maintenance of the Training Material repository of the Galaxy Training Network
- Management of Galaxy Training community
- Member of the training group of de.NBI, the German Network for Bioinformatics infrastructure
- Development of command-line tools, tools for Galaxy, packages for BioConda
- Teaching to computer science master students

Post-doctoral researcher

Clermont-Ferrand, France

Jan. 2015 - Jun. 2016

CIDAM, Université d'Auvergne

- Development of metagenomics tools for Galaxy
- Data analysis of metagenomics data
- Teaching to bioinformatics undergrad students

PhD student & researcher

Lyon, France

Oct. 2011 - Nov. 2014

Institut National des Sciences Appliquées

- In silico experimental evolution
- Comparative genomics with phylogeny
- Teaching to bioinformatics undergrad students

Extracurricular Activity

Mozilla Open Leader

Mozilla Foundation Sep. 2018 - Today

- Mentoring
- 14 week leadership development program

March 14, 2019 Bérénice Batut · Curriculum Vitae

Mentor

THE CARPENTRIES Sep. 2018 - Today

· Mentoring on community building and material development

Member

TRAINING GROUP (SIG3) OF THE GERMAN NETWORK FOR BIOINFORMATICS INFRASTRUCTURE

Dec. 2016 - Today

Instructor

THE CARPENTRIES March. 2016 - Today

- Training for foundational computational and data science skills to researchers
- Unix Shell, Version Control with Git, Plotting and Programming with Python and R

Training & Teaching

The complete list of publications can be found: http://research.bebatut.fr/training

2019

Sep. 18	Galaxy data analysis (introduction, ChiP-seq, RNA-seq), 5 days workshop	Freiburg, Germany
Jan. 22	Galaxy RNA-seq data analysis, 2 days workshop	Belgrade, Serbia
2018		
Oct. 18 -	Introduction to data-driven life science, Lectures and exercises to computer science master students	Freiburg, Germany

Feb. 19

Dec. 18 Galaxy RNA-seq data analysis, 2 days workshop

Sep. 18 Galaxy data analysis (introduction, ChiP-seq, RNA-seq), 5 days workshop

Jun. 18 Galaxy Introduction and Galaxy for Training, 2 2.5 h workshops at the Galaxy Community Conference

May. 18 Galaxy data analysis (introduction metagenomics), 2 days workshop

Apr. 18 Galaxy data analysis (introduction metagenomics), 2 days workshop

Cape Town, South

Apr. 18 **Galaxy data analysis (introduction, metagenomics),** 2 days workshop

Cape Town, South Africa

Africa

Africa

Africa

Africa

Cape Town, South Africa

Africa

**Africa

Feb. 18 Galaxy data analysis (introduction, ChiP-seq, RNA-seq), 5 days workshop

Freiburg, Germany

Publications

The complete list of publications is available online: http://research.bebatut.fr/publications

The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2018 update.

Batut B et al Afgan E

Nucleic acids research (May 2018). 2018

ASaiM: a Galaxy-based framework to analyze microbiota data

Batut B al

GigaScience (May 2018). Oxford University Press (OUP), 2018

Community-Driven Data Analysis Training for Biology.

Batut B al

Cell systems (June 2018). 2018

The RNA workbench: Best practices for RNA and high-throughput sequencing bioinformatics in Galaxy Grüning B A, J. Fallmann, D. Yusuf, S. Will, A. Erxleben, F. Eggenhofer, T. Houwaart, Batut B al

Nucleic Acids Research 45.W1 (2017) W560-W566. 2017

ENASearch: A Python library for interacting with ENA's API

Batut B, Gruening BA

The Journal of Open Source Software (2017). 2017

Environmental pollutant Benzo[a]Pyrene impacts the volatile metabolome and transcriptome of the human gut microbiota Defois C, Ratel J, Denis S, Batut B al

Frontiers in Microbiology 8.AUG (2017). 2017

Four simple recommendations to encourage best practices in research software

Jiménez R J, Kuzak M, Alhamdoosh M, Barker M, Batut B al

F1000Research 6 (June 2017) p. 876. F1000 Research, Ltd., 2017